

# Estrogen Regulation of Epididymal Sperm Maturation

Michelle Krisfalusi

*Gamete Biology Section  
Laboratory of Reproductive and Developmental Toxicology  
National Institutes of Environmental Health Sciences  
National Institutes of Health*

# Background

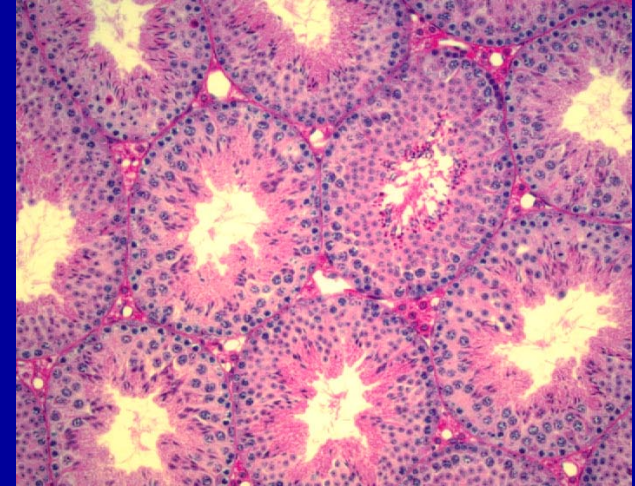
- The target tissues of estrogens and the specific functions of estrogen receptors in male reproductive function are unknown.
- Estrogen receptor  $\alpha$  (ER $\alpha$ ) and  $\beta$  (ER $\beta$ ) are expressed in a wide variety of cell-types in the male reproductive tract.
- The highest levels of ER $\alpha$  expression are found in the epithelial cells of the efferent ducts and caput region of the epididymis in rodents.
- $\alpha$ ERKO mice are infertile, while  $\beta$ ERKO mice are fertile.
- Infertility of  $\alpha$ ERKO males is due to disruption of ER action within somatic cells and not germ cells (Mahato et al., 2000).

# $\alpha$ ERKO Male Mice Are Infertile

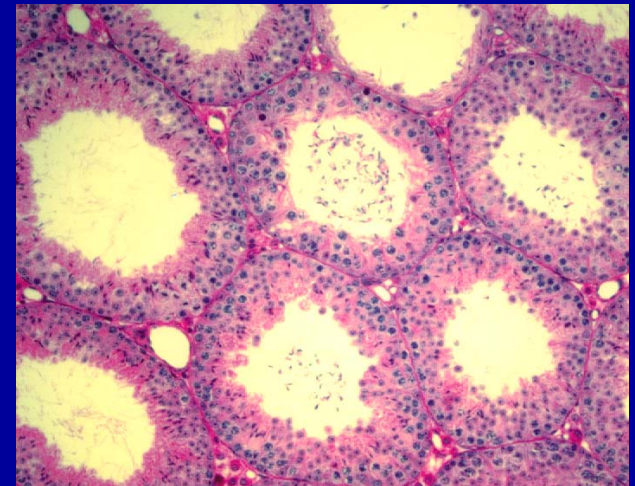
## ER $\alpha$ -/- phenotype:

- Altered testicular morphology
- Dilated rete testis
- Reduced mating frequency
- Reduced epididymal sperm count
- Reduced sperm motility
- Sperm fail to fertilize eggs *in vitro*

Wild Type

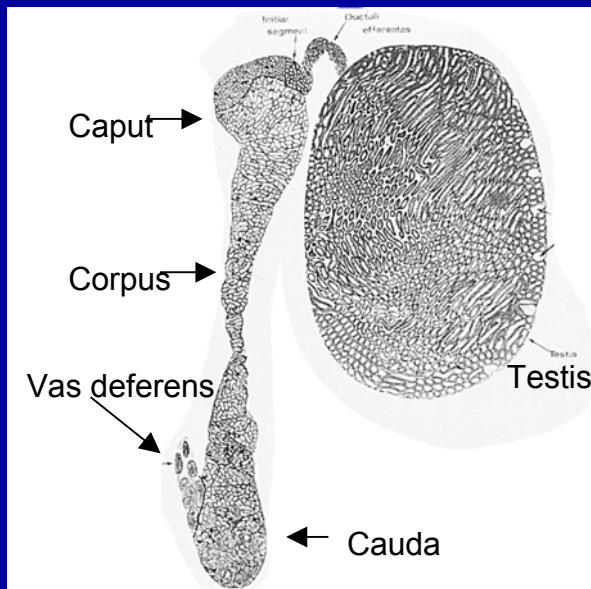


$\alpha$ ERKO



# Epididymis

- Functions as a storage site and in the final maturation of sperm.
- ER $\alpha$  is present in epithelial cells with highest levels in the caput.
- Epithelial cells modify the luminal fluid and alter biochemical and physiological properties of sperm.
- Sperm maturation and transport are reported to be influenced by estrogens, but few studies have been done.



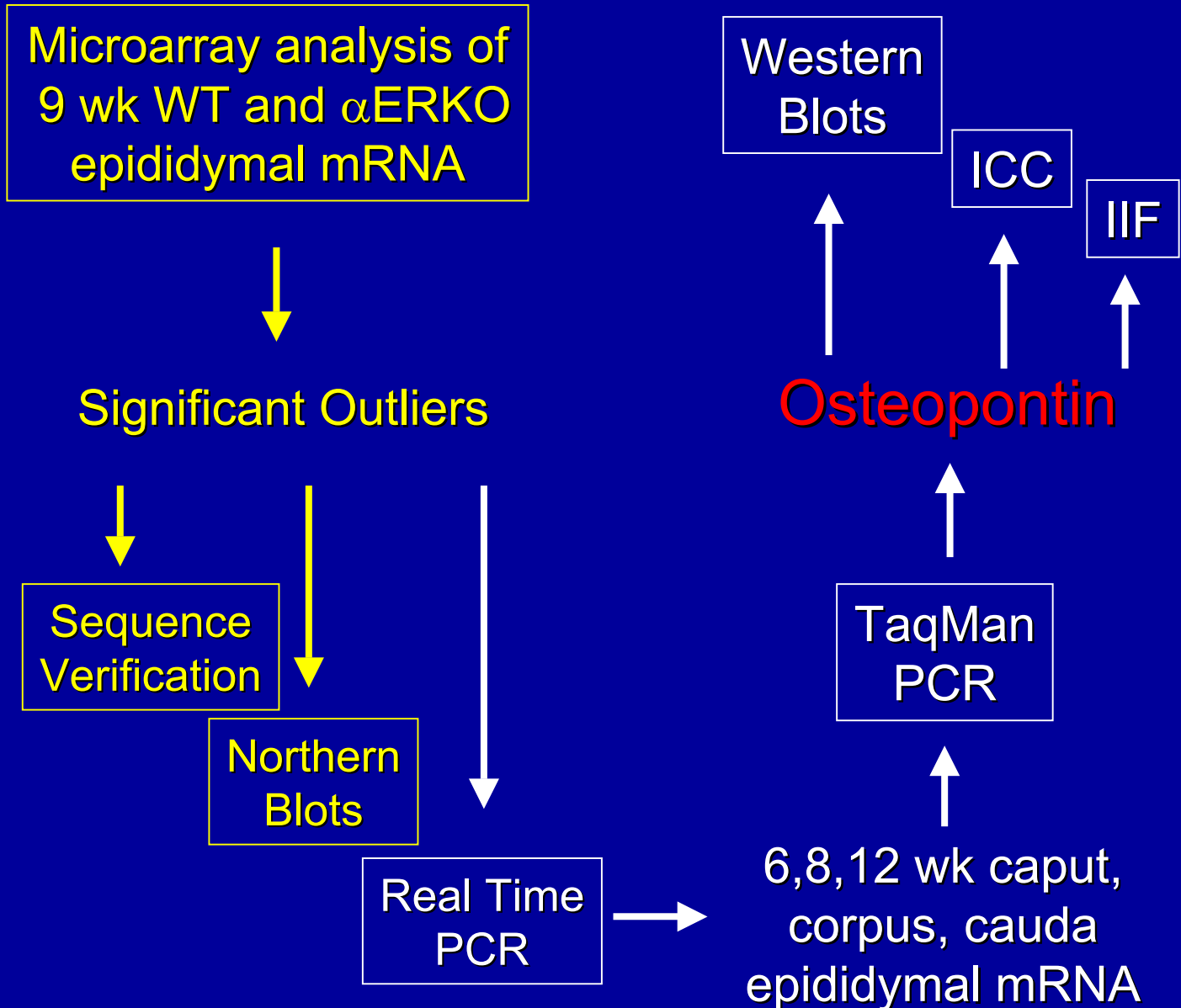
Cauda sperm of  $\alpha$ ERKOs exhibit low motility and fail to fertilize eggs suggesting that sperm of  $\alpha$ ERKO mice fail to undergo epididymal maturation.

Rationale: Sperm acquire motility and the ability to fertilize eggs as they transit the epididymis. Epididymal sperm of  $\alpha$ ERKO mice fail to undergo these changes. Therefore, it is suggested that ER $\alpha$  and estrogens play a fundamental role in male reproduction by regulating critical aspects of epididymal sperm maturation.

Hypothesis: Estrogen regulates the synthesis of proteins by epididymal cells required for sperm to undergo the physiological changes and surface modifications necessary to become capable of fertilization.

Goal: To identify estrogen-regulated proteins that are involved in the epididymal maturation of sperm and determine their roles in this process.

# Experimental Design



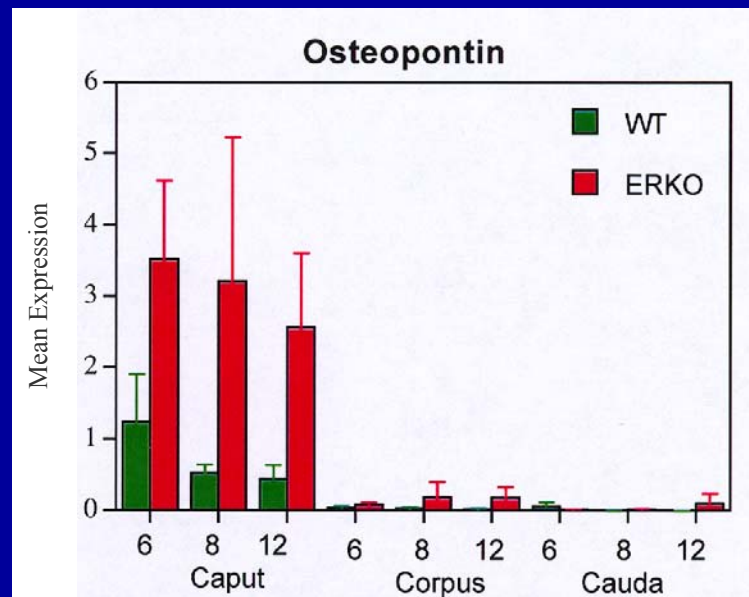
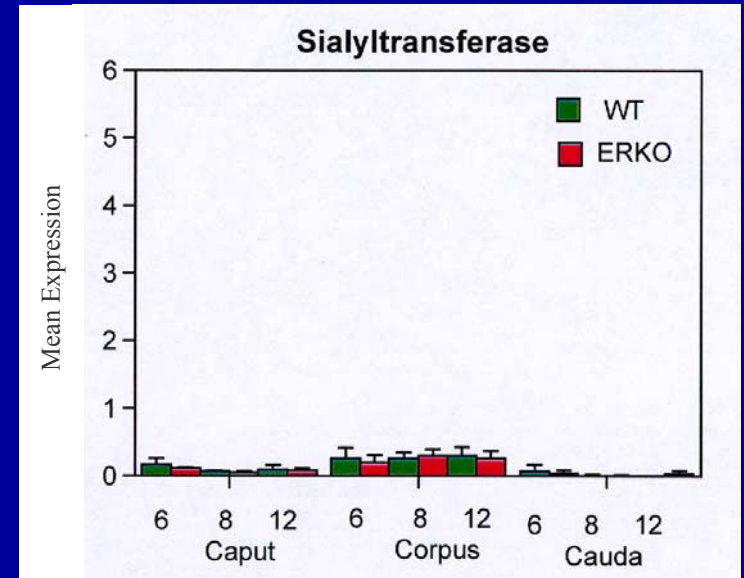
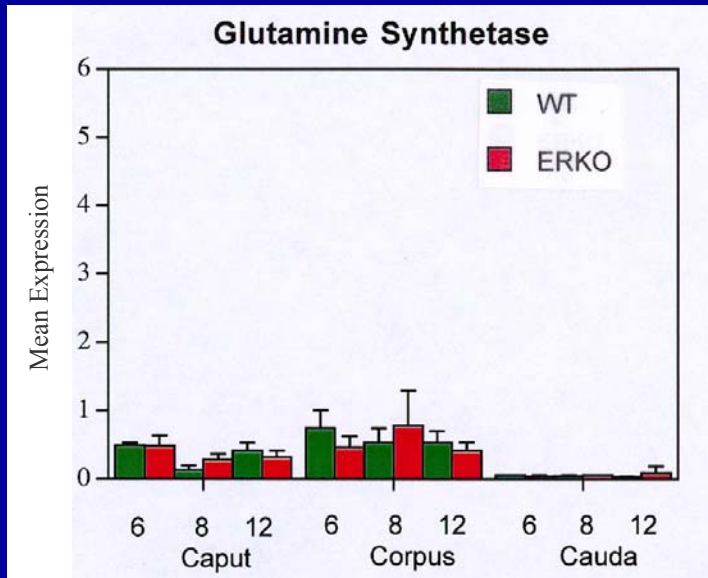
Clone Identification	Expression Levels			Sequence
	Microarray	Northern	Real Time PCR	
Hemaglobin alpha	2.1x ERKO	.9x ERKO		NO sequence
EST	1.7x ERKO	2.3x ERKO		YES
EST	1.6x ERKO			NO sequence
EST	1.6x ERKO	1.7x ERKO		NO
Osteopontin	1.5x ERKO	2.1x ERKO	4.3x ERKO	YES
Thymopoietin alpha	1.5x ERKO	1.8x ERKO		YES
EST	1.4x ERKO			NO sequence
Mus Xlr4 mRNA	1.4x ERKO	2.3x ERKO		YES
mRNA for ribonucleoprotein H1	.7x ERKO	1.2x ERKO		YES
Solute carrier family 31	.7x ERKO	.8x ERKO		NO
Heat shock protein cognate 70	.7x ERKO			YES
EST	.7x ERKO			YES
Phosphodiesterase	.7x ERKO	.9x ERKO		YES
Calcium binding protein	.7x ERKO			NO sequence
EST	.6x ERKO	.9x ERKO		NO
Glutamine synthetase	.6x ERKO	.9x ERKO		YES
EST	.6x ERKO	.7x ERKO		YES
EST	.6x ERKO	.7x ERKO		YES
Sialyltransferase	.6x ERKO	.4x ERKO	.6x ERKO	YES
SSeCKS, testis specific gene A1	.6x ERKO	.4x ERKO	.5x ERKO	YES
ATP-binding cassette	.6x ERKO	.8x ERKO		YES
Metallothionein 2	.5x ERKO	.7x ERKO		NO



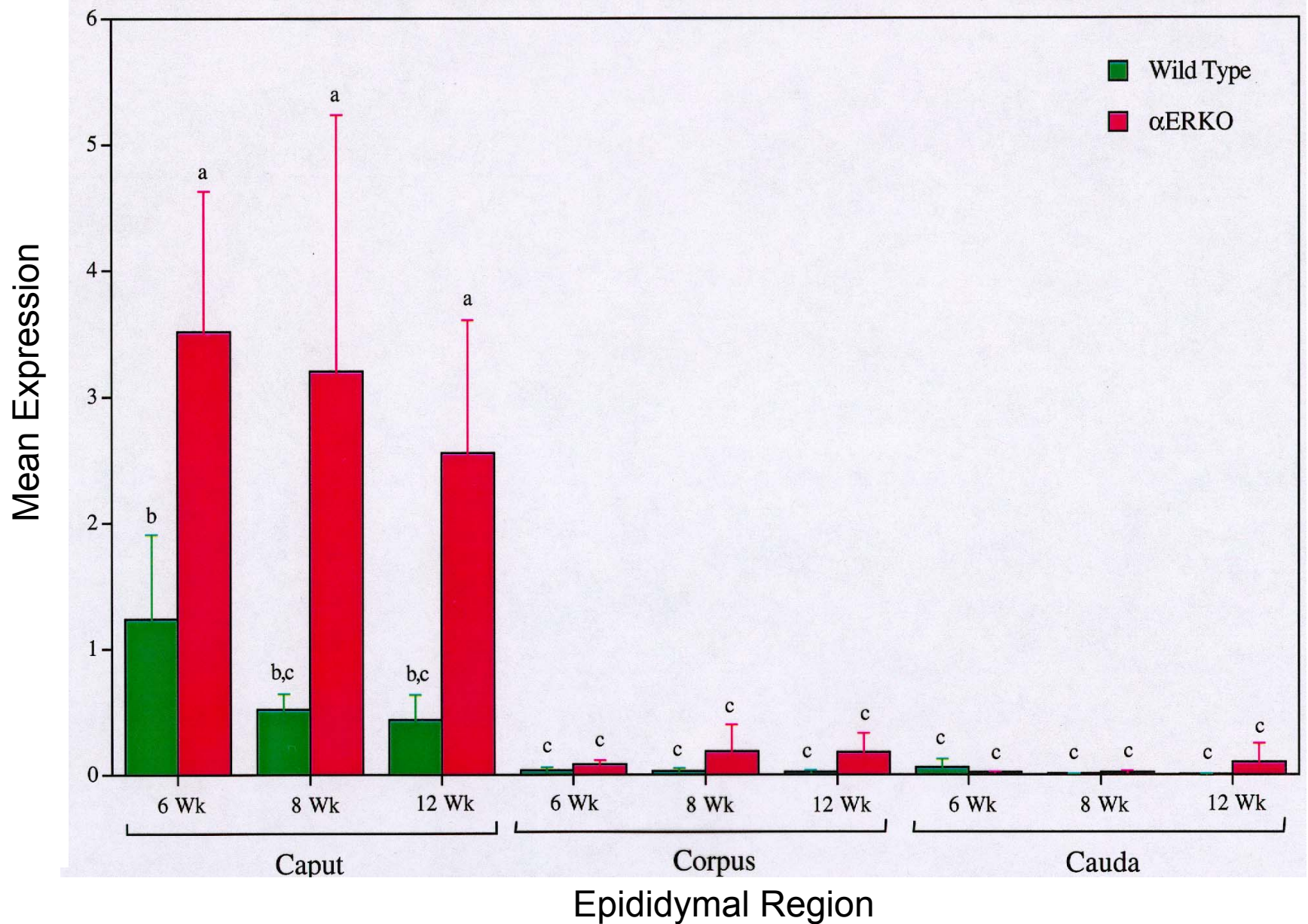
Objective: To measure differential expression of selected genes by collecting caput, corpus, and cauda epididymides from 6, 8, and 12 week WT and  $\alpha$ ERKO males for real time PCR analysis.

Hypothesis: The caput, corpus and caudal regions of the epididymis will show differential gene expression and the results gathered from microarray analysis will be enhanced following real time PCR analysis.

# TaqMan PCR



# Osteopontin Gene Expression

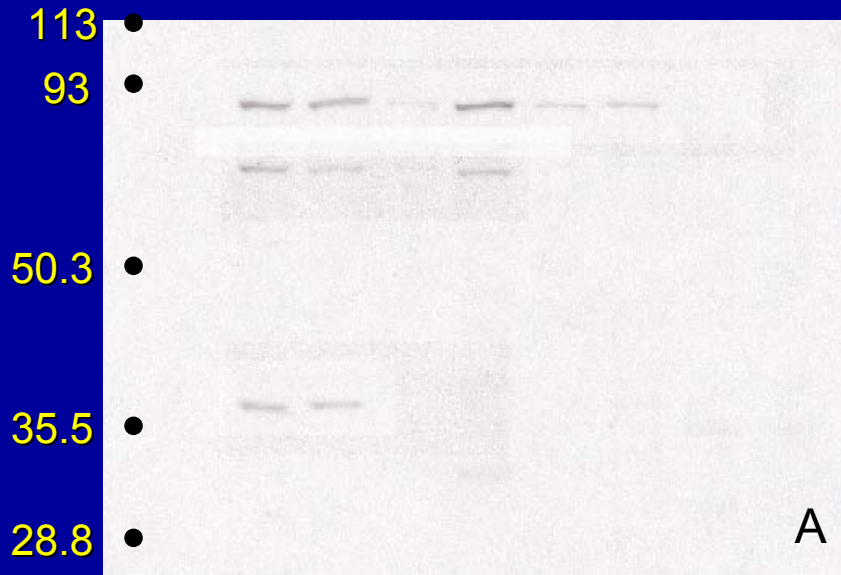


# Osteopontin

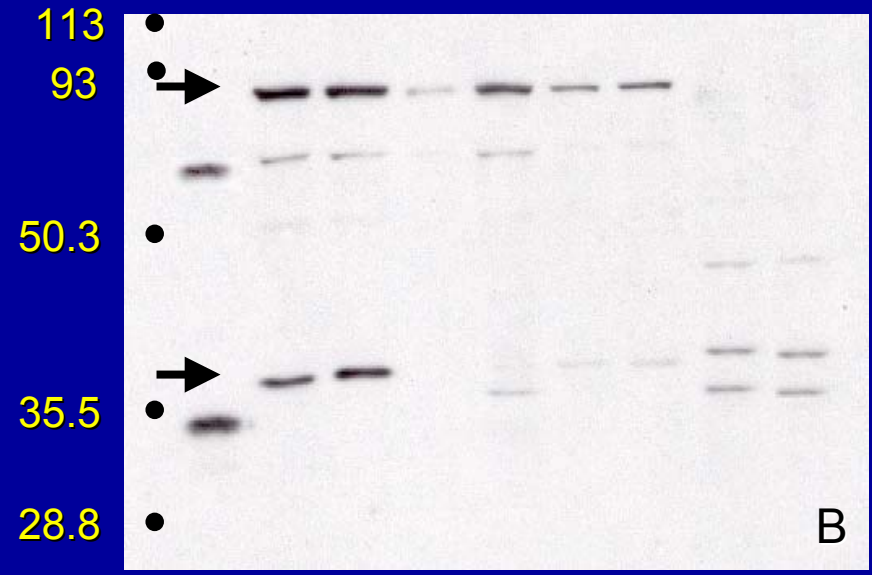
- Originally isolated from bone matrix and is the principal phosphorylated glycoprotein of bone.
- It is expressed in a wide variety of mammalian tissues and biological fluids.
- Highly acidic, calcium binding, secreted phosphorylated glycoprotein.
- MW has been reported to range from 31-75 KDa due to post-translational modification and splice variants.
- Single copy gene with two half-EREs.

# Western Blot

Recombinant OPN  
 WT Caput Epididymis  
 ERKO Caput Epididymis  
 WT Cauda Epididymis  
 ERKO Cauda Epididymis  
 WT Caput Sperm  
 ERKO Caput Sperm  
 WT Cauda Sperm  
 ERKO Cauda Sperm



Recombinant OPN  
 WT Caput Epididymis  
 ERKO Caput Epididymis  
 WT Cauda Epididymis  
 ERKO Cauda Epididymis  
 WT Caput Sperm  
 ERKO Caput Sperm  
 WT Cauda Sperm  
 ERKO Cauda Sperm



# Osteopontin and Male Reproduction

- OPN has been identified in rat epididymis and on rat and bull epididymal sperm.
- It has been reported that the immunoreactivity of OPN on the sperm head in rats disappears as sperm transit the epididymis (Sitteri et al., 1995).
- It has been suggested that the presence of OPN, a calcium-binding protein, in the epididymis may serve as a “decapacitation” factor to prevent premature activation of epididymal sperm motility or fertilizing ability (Sitteri et al., 1995).
- Hypothesis: over-expression of OPN in  $\alpha$ ERKO males may lead to the reduced sperm motility and the inability to fertilize eggs.



# Conclusions

- 22 genes were found to be differentially expressed in 9 wk WT and  $\alpha$ ERKO epididymides following microarray analysis.
- Based on validated microarray results and potential relevance to epididymal function and estrogen regulation, 3 genes were chosen for further exploration using TaqMan PCR.
- TaqMan PCR indicated that osteopontin was up-regulated in the caput region of the epididymis in  $\alpha$ ERKOs compared to WT.
- Western blot analysis has indicated OPN protein is expressed in the caput and cauda regions of WT and  $\alpha$ ERKO animals with an increase in expression in the cauda region of  $\alpha$ ERKOs compared to WT animals.

# Future Studies

1. Complete Immunohistochemistry to localize OPN protein expression in the epididymis.
2. Examine OPN protein expression on epididymal sperm by using indirect immunofluorescence.
3. Microarray analysis of WT and  $\alpha$ ERKO caput epididymis to determine if additional differentially expressed genes can be determined.



# Acknowledgements

Mitch Eddy  
Dipak Mahato  
Ken Korach  
Vicki Walker  
Microarray center  
Cindy Afshari  
Rick Paules  
Chris Miller  
Nigel Walker

